

FOOT 507650

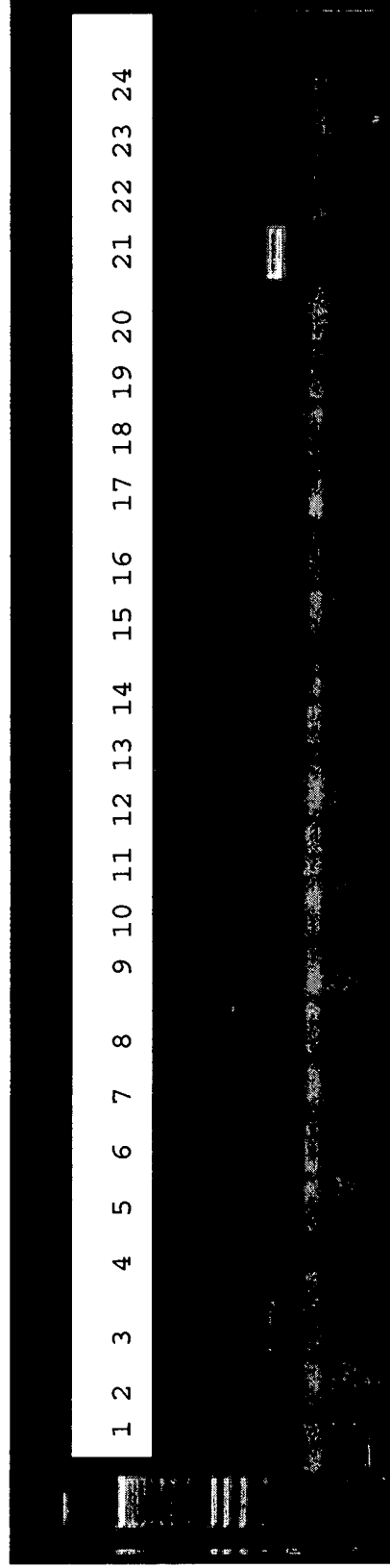


Figure 1

Mouse	MGGCEVREFLLQFGFFLPLLTAWTGDCSHVSN-QVVLLDTTVMGELGWKTYPLNGWDAI	59
Human	MGGCEVREFLLQFGFFLPLLTAWPGDCSHVSNQVVLLDTTTLGELGWKTYPLNGWDAI	60

Mouse	TEMDEHNRPIHTYQVCNVMENQNNWLRNWNISRDAAQKIYVEMKFTLRDCNSIPWVLGT	119
Human	TEMDEHNRPIHTYQVCNVMENQNNWLRNWNISRDAAQKIYVEMKFTLRDCNSIPWVLGT	120

Mouse	CKETFNLYIESDESHGTFKFKPSQYIKIDTIAADESFTQMDLGDRIKLNTEIREVGPIE	179
Human	CKETFNLYMESDESHGTFKFKPNQYTKIDTIAADESFTQMDLGDRIKLNTEIREVGPIE	180

Mouse	RKGFYLAQDQIGACIALVSVRVFYKKCPFTVRSLAMFPDTIPRVDSSSLVEVRGSCVKSA	239
Human	RKGFYLAQDQIGACIALVSVRVFYKKCPFTVRNLMAMPDTIPRVDSSSLVEVRGSCVKSA	240

Mouse	EERDTPKLYCGADGDWLVPLGRICISTGYEEIEGSCHACRPGFYKAFAGNTKCSKCPPHS	299
Human	EERDTPKLYCGADGDWLVPLGRICISTGYEEIEGSCHACRPGFYKAFAGNTKCSKCPPHS	300

Mouse	STYVEATSVCHCEKGYFRAEKDPPSMACRPPSAPRNVAFNINETALILEWSPPSDTGGR	359
Human	LTMEATSVQCCEKGYFRAEKDPPSMACRPPSAPRNVAFNINETALILEWSPPSDTGGR	360
	**	
Mouse	KDLTYSVICKKCGLDTTQCEDCGGGLRFIPRHTGLINNSVVVLDVFSHVNYTFEIAMNG	419
Human	KDLTYSVICKKCGLDTSQCEDCGGGLRFIPRHTGLINNSVIVLDVFSHVNYTFEIAMNG	420

Mouse	VSELSISPKPFTAITVTTDHDAPSLIGMMRKDWASQNSLALSQWAPAFSNGAILDYETKY	479
Human	VSELSFSPKPFATAITVTTDQDAPSLIGVVRKDWASQNSIALSQWAPAFSNGAILDYETKY	480

Mouse	YEKEHEQLTYSSTRSKAPSVIITGLKPATYIFHIRVRTATGYSQKFEFETGDETS	539
Human	YEKEHEQLTYSSTRSKAPSVIITGLKPATYIFHIRVRTATGYSQKFEFETGDETS	540

Mouse	MAAEQGGQILVIATAAVGGFTLLVILTLFFLITGRCQWYIKAKMKSEKRRTHLQNGHLRF	599
Human	MAAEQGGQILVIATAAVGGFTLLVILTLFFLITGRCQWYIKAKMKSEKRRNHLQNGHLRF	600

Mouse	PGIKTYIDPDYEDPSLAVHEFAKEIDPSRIRIERVIGAGEFGEVCSGRLKTPGKREIPV	659
Human	PGIKTYIDPDYEDPSLAVHEFAKEIDPSRIRIERVIGAGEFGEVCSGRLKTPGKREIPV	660

Mouse	AIKTLKGGHMDRQRDFLEASIMGQFDHPNIIRLEGVVTKRSPFAIGVEAFPCPSFLRAG	719
Human	AIKTLKGGHMDRQRDFLEASIMGQFDHPNIIRLEGVVTKRSPFAIGVEAFPCPSFLRAG	720

Mouse	FLNGIQAPHPVTAGGSLPPRIAPGRPVMIIVVEYMENGLSDSLRKHGDHFTVIQLVGMRL	779
Human	FLNSIQAPHPVPGGSLPPRIAPGRPVMIIVVEYMENGLSDSLRKHGDHFTVIQLVGMRL	780

Mouse	GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFLSRVLEDDPEAYTTTGKGIPI	839
Human	GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFLSRVLEDDPEAYTTTGKGIPI	840

Mouse	RWTAPEAIAIRKFSASDAWSYGIVMWEVMSYGERPYWEMSNQDVILSIEEGYRLPAPMG	899
Human	RWTAPEAIAIRKFSASDAWSYGIVMWEVMSYGERPYWEMSNQDVILSIEEGYRLPAPMG	900

Mouse	CPPSLQQLMLHCWQKERNHRPKFTDIVSFLDKLIKNPSALHTLVEDILVMPESPGDVPEY	959
Human	CPASLHQLMLHCWQKERNHRPKFTDIVSFLDKLIKNPSALHTLVEDILVMPESPGEVPEY	960
	**	
Mouse	PLFVTVGDWLDSEIKMGQYKSNFMAAGFTTFLISRMSIDDIRRIGVILIGHQRRIVSSIQ	1019
Human	PLFVTVGDWLDSEIKMGQYKSNFMAAGFTTFLISRMSIDDIRRIGVILIGHQRRIVSSIQ	1020

Mouse	TLRLHMMHIQEKGFHV	1035
Human	TLRLHMMHIQEKGFHV	1036

FIGURE 2

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Human: 903 aggggttcttgccatgcttgacagaccaggattctataaagcttttgctgggaacacaaaat 962
 Mouse: 1099 aggggttcttgccatgcttgacagaccaggattctacaagcatttgctgggaacacaaaat 1158

Human: 963 gttctaaatgtcctccacacagtttaacatacatggaagcaacttctgtctgtcagtgtg 1022
 Mouse: 1159 gttccaaatgccctccacacagctcaacctacgtggaagcaacgtcagtctgtcattgtg 1218

Human: 1023 aaaaggggtattttccgagctgaaaaagaccaccttctatggcatgtaccaggccacctt 1082
 Mouse: 1219 aaaaggggtacttccgggcagaaaaagaccaccttctatggcatgcactagaccacctt 1278

Human: 1083 cagctcctaggaatgtgggttttaacatcaatgaaacagcccttattttggaatggagcc 1142
 Mouse: 1279 cagctcctagaaatgtggcttttaacatcaatgaaacagcccttattttggaatggagcc 1338

Human: 1143 caccaagtgcacaggaggagaaaagatctcacatacagtgaatctgtaagaaatgtg 1202
 Mouse: 1339 caccagtgacacaggaggagaaaagatctcacatacagtgaatctgtaagaaatgtg 1398

Human: 1203 gcttagacaccagccagtgaggactgtggtggaggactccgcttcatcccaagacata 1262
 Mouse: 1399 gtttagacactacccagtgaggactgtggtggaggactccgcttcatcccaagacaca 1458

Human: 1263 caggcctgatcaacaattccgtgatagtacttgactttgtgtctcacgtgaattacacct 1322
 Mouse: 1459 ctggactgatcaacaattctgtgtagtactggactttgtgtctcacgtcaattataacct 1518

Human: 1323 ttgaaatagaagcaatgaatggagtttctgagttgagtttttctcccaagccattcacag 1382
 Mouse: 1519 ttgaaatagaagccatgaatggagtttctgagttgagcatctctcccaagccattcacag 1578

Human: 1383 ctattacagtgaccacggatcaagatgcaccttccctgatagggtgtgtaaggaaggact 1442
 Mouse: 1579 ctattacagtgactacagatcacgatgcaccttctctgattggtatgatgagaaaggact 1638

Human: 1443 gggcatcccaaaatagcattgccctatcatggcaagcacctgctttttccaatggagcca 1502
 Mouse: 1639 gggcatctcagaacagccttgctctatcgtggcaagcacctgcattttccaatggagcta 1698

Human: 1503 ttctggactacgagatcaagtactatgagaaagaacatgagcagctgacctactcttcca 1562
 Mouse: 1699 ttctggactatgagaccaagtactacgagaaagagcatgagcagctcacctattcttcca 1758

Human: 1563 caaggtccaaagccccagtgatcatcatcacaggtcttaagccagccaccaaataatgtat 1622
 Mouse: 1759 cgagggtccaaagccccagcgtcatcgtcacgggctcaagcccgccaccagctacatat 1818

Human: 1623 ttcacatccgagtgagaactgcgacaggatacagtggtacagtcagaaatttgaatttg 1682
 Mouse: 1819 ttcacatccgagtgaggacggcgacaggctacagtggtacagtcagaaatttgaatttg 1878

Human: 1683 aaacaggagatgaaacttctgacatggcagcagaaacaggacagattctcgtgatagcca 1742
 Mouse: 1879 aaacaggagatgaaacttctgacatggcggcagaaacaggacagattctcgtgatagcca 1938

Human: 3483 gggaagtgttcacggacttaacctaataaaatttatcc--aggtggggcttccttagtga 3540
|||||
Mouse: 3678 gggaagtgttcacggacttaacctaataaggaattgtccttaagtgtggcatctcagaga 3737

Human: 3541 tgtatgtagagtgtgatgtagagaaagaactagttgacctttctttcatgttttgt 3600
|| |||
Mouse: 3738 cgtgtgtggagtttgatgatagataagaagaaaatagctgatcattcttccatgttttgt 3797

Human: 3601 gatcaagtagct 3612
||||
Mouse: 3798 gatcgagtagct 3809

Human: 3601 gatcaagtagct 3612

Figure 4

KSE132	MGGCEVREFFLLQFGFFLPLLTAWPGDCSHVSNNQVLLDTTTTLGELGWKTYPLNGWDAI	60
XM_035574	-----	
KSE132	TEMDEHNRPIHTYQVCNVMEPNQNNWLRTNWSRDAQAQKIYVEMKFTLRDCNSIPVWLGT	120
XM_035574	-----	
KSE132	CKETFNLFYMESDESHGIKFKNPYTKIDTIAADESFTQMDLGDRILKLNTIEIREVGPIE	180
XM_035574	-----	
KSE132	RKGfYLAFQDIGACIALVSVRVFYKKCPFTVRNLAMFPDTIPRVDSSSLVEVRGSCVKSA	240
XM_035574	-----	
KSE132	EERDTPKLYCGADGDWLVLPLGRICISTGYEEIEGSCHACRPGFYKAFAGTKCSKCPPHS	300
XM_035574	-----	
KSE132	LTYMEATSVQCQCEKGYFRAEKDPSPMACTRPSPAPRNVVFNINETALILEWSPPSDTGGR	360
XM_035574	-----	
KSE132	KDLTVSICKKCLDTSQCEDCGGLRFIPRHGTGLINNSVIVLDFVSHVNYTFEIAMNG	420
XM_035574	-----	
KSE132	VSELFSFSPKPFTAITVTTDQAPSLIGVVRKDWASQNSIALSWQAPAFSNGAILDYEIKY	480
XM_035574	-----	
KSE132	YEKEHEQLTYSSTRSKAPSVIITGLKPKATKYVFHIRVRTATGYSYGYSQKFEFETGDESD	540
XM_035574	-----MKXXXXXXXXXXXXXXXXXXXXXSD :: **	26
KSE132	MAAEQGQILVIATAAVGGFTLLVILTFLFLITGRCQWYIKAKMKSEEKRRLHLQNGHLRF	600
XM_035574	MAAEQGQILVIATAAVGGFTLLVILTFLFLITGRCQWYIKAKMKSEEKRRLHLQNGHLRF *****	86
KSE132	PGIKTYIDPDITYEDPSLAVHEFAKEIDPSRIERIIVIGAGEFGEVCSGRLKTPGKREIPV	660
XM_035574	PGIKTYIDPDITYEDPSLAVHEFAKEIDPSRIERIIVIGAGEFGEVCSGRLKTPGKREIPV *****	146
KSE132	AIKTLKGGHMDRQRDFLREASIMGQFDHPNIIRLEGVVTKRSFPAIGVEAFCPSFLRAG	720
XM_035574	AIKTLKGGHMDRQRDFLREASIMGQFDHPNIIRLEGVVTKRSFPAIGVEAFCPSFLRAG *****	206
KSE132	FLNSIQAPHVPVGGGSLPPRIAPGRPVMIVVEYMENGLSDSFLRKHDGHFTVIQLVGMLR	780
XM_035574	FLNSIQAPHVPVGGGSLPPRIAPGRPVMIVVEYMENGLSDSFLRKHDGHFTVIQLVGMLR *****	266
KSE132	GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFGLSRVLEDDPEAAYYTTTGKGIPI	840
XM_035574	GIASGMKYLSDMGYVHRDLAARNILVNSNLVCKVSDFGLSRVLEDDPEAAYYTTTDD----	321
KSE132	RWTAPEAIAYRKFFSSASDAWSYGIVMWEVMSYGERPYWEMSNDQVILSIEEGYRLPAPMG	900
XM_035574	-----LFQTLLTNLCYSA----- *.:. .*.	334
KSE132	CPASLHQLMLHCWQKERNHRPKFTDIVSFLDKLIRNPALHTLVIEDILVMPESPGEVPEY	960
XM_035574	-----	
KSE132	PLFVTVDGWLDSIKMGQYKNNFVAAGFTTFDLISRMSSIDIRRIGVILIGHQRRIVSSIQ	1020
XM_035574	-----	
KSE132	TLRLHMMHIQEKGPHV	1036
XM_035574	-----	